

Nucleotide Sequences pXP10 (SEQ ID No: 111)

1 GACGAAGGG CCTCGTATA CGCCATTTAT TATAGGTTAA TGTCTATGATA  
CTGCTTCCC GGAGCACTAT CGGGATAAAA ATATCCAATT ACAGTACTAT  
51 ATAATGGTT CCTAGACGTC AGGTGGCACT TTTCGGGAA ATGTGCGCG  
TATTACCAA GAATCTGAG TCCACCGTGA AAAGCCCCCT TACACGGGCC  
101 AACCCCTATT TGTTTATTTT TCTAAATACA TTCAAAATATG TATCCGCTCA  
TTGGGGATAA ACAAAATAAA AGATTTATGT AAGTTTATAC ATAGGCAGT  
151 TGAGACAATA ACCCTGATAA ATGCTTCAAT AATATTGAAA AAGGAAGAGT  
ACTCTGTTAT TGGGACTATT TAGAAGTTA TTATAACTTT TTCCCTCTCA  
201 ATGAGTATTAC AACATTCCG TGTGCCCTT ATTCCCTTT TTGCGGATT  
TACTGATAAG TTGTAAGGC ACAGCGGGAA TAAGGGAAAA AACGCCGTA  
251 TTAGGTTTGCT GTTTTGTGTC ACCCGAACAA GCTGTGAAAG GTAAAAGATG  
AACGGAAGGA CAAAACAGG TGGGTCTTT CGACCACTT CATTTCTAC  
301 CTGAAGATCA GTTGGGTGCT CGAGTGGGTT ACATCGAACT GGATCTCAAC  
GACTCTGAGT CAACCCACGA GCTCACCCAA TGATCCTTGA CCTAGAGTTG  
351 AGCGGTAAGA TCTTGTAGAG TTTTGGCCC GAAGAACGTT TTCAATGAT  
TGGCATTCT AGGAACCTCT AAAAGCGGGG CTTCTTGAA AAGGTTACTA  
401 GAGCACTTT AAAGTCTGC TATGTTGGCC GGTTATTTAC CGTATTGACG  
CTCGTGAAA TTTCAAGACG ATACACCGG CCATAATAGG GCATAACTGC  
451 CCGGGCAAGA GCAACTCGGT CGCCGCATAC ACTATTCTCA GAATGACTTG  
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501 GTTGTAGTACT CACAGTCAC AGAAAAGCAT CTTACGGATG GCATGACAGT  
CAACTCATGA GTGGTCAGTG TCTTTCTGA GAATGCCAT CGTACTGTCA  
551 AAGAGAATTA TGCAGTGTG CGATAAACCT GAGTGATAAC ACTGCGGCCA  
TTCTCTTAAT CGTCAGCACG CGTATTGTTA CTCACTATTG TGACGCCGT  
601 ACTTACTTCT GACAACGATC GGAGGACCGA AGGAGCTAAC CGCTTTTTG  
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651 CACAACATGG GGGATCATGT AACTCGCTT GATCGTTGGG AACCGGAGCT  
GTGTTGTACC CCCTAGTACA TTGAGCGGGAA CTAGCAACCC TTGGCCTCGA  
701 GAATGAAGCC ATACCAACAG ACAGCGCTGA CACCAACGATG CCTGTAGCAA  
TTCTACTTCG TATGTTGTC TGCTGCACT GTGGTGCTAC GGACATCGT  
751 TGGCAACAC GTTGCGBAAA CTATTAACCTG GCGAACTACT TACTCTAGCT  
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801 TCCGGCACR AATTAAATAGA CTGGATGGAG CGGGATAAAG TTGAGGACC  
AGGGCGTTG TAAATTATCT GACCTACCTC CGCCTATTTC AACGTCTTGG  
851 ACTTCTGCGC TCGGGCCCTC CGGGCGCTG GTTTATTGCT GATAAATCTG  
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901 GAGCCGGTGA CGCTGGGTCT CGCGGTATCA TTGAGCACT GGGGGCAGAT  
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951 GGTAAAGCCCT CCCGTATCGT AGTTATCTAC ACGACGGGG A GTCAGGCAAC  
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1001 TATGGATGAA CGAAATAGAC AGATCGCTGA GATAGGTGCC TCACGTGATTA  
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1301 TTGTTTGCGC GATCAAGAGC TACCAACTCT TTTTCCGAAG GTAATGCGT  
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1351 TCAGCAGAGC CGACAGATCCA AATACTGTCC TTCTAGTGTAA CGCGTAGTTA

-Replacement Sheet-

10/43

1401 AGTCGTCTCG CGTCTATGGT TTATGACAGG AAGATCACAT CGGCATCAAT  
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CCGGTGGTGA AGTTCTTGTAG ACATCGTGG GGATGTATGG AGCAGAGCGA  
1451 AATCTGTGA CCAGTGGCTG CTGCCAGTGG CGATAAGTCG TGTCTTACCG  
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1501 GTTGGACTC AAGACGATAG TTACCGATA AGGCCACGG GTCGGCTGA  
CCAACCTGAG TTCTGCTATC AATGGCCTAT TCCGGTCCG CAGCCCGACT  
1551 CGGGGGGGTGT CGTGCATACA GCCAGCTTG GAGCGAACGA CCTACACCGA  
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1601 ACTGAGATAC CTACAGCGTG AGCTATGAGA AAGCGCCACG CTTCGGAAAG  
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1651 GGAGAAAGGC GGACAGGTAT CGGTAAGCG GCAGGGTCCG AACAGGAGAG  
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1701 CGCACGAGGG AGCTTCCAGG GGGAAACGCC TGGTATCTT ATAGTCCCTG  
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1751 CGGGTTTCCG CACCTCTGAC TTGAGCGTGC ATTGTTGTGA TGCTGTCAG  
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1801 GGGGGCGGGAG CCTATGGAAA AACGCCAGAA AGCGGCCCTT TTAGGGTTC  
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1951 CGCGCAGCGG AACACGGCGG CGCACCGAGT CAGTGAGCGGA GGAAGCGGAA  
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2001 GAGCGCCCAA TACGCAAACCC GCCTCTCCCC CGCGCTTGGC CGATTCTATTA  
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2051 ATGCACTGG CACGACAGGT TCCCGACTG GAAACGGGGC AGTGGCCGA  
TACGTGACCC GTGCTGTCCA AAGGGCTGAC CTTTGGCCCG TCACTCGCT  
2101 ACGCAATTAA TGTGAGTGTG CTCACTTATT AGGCACCCCCA GGCTTTACAC  
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2151 TTAATGCTTC CGGCTCGTAT GTTGTGTGGA ATTGTGAGGG GATAACAATT  
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2201 TCACACAGGA AACACGTATG ACCATGATTA CGCAAGCTT TGGAGCCTT  
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2251 TTTGGAGA TTTCACAGT GAAAATTAAT TTATTCGCAA TTCCCTTTAGT  
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2301 TGTTCTTTC TATGGGGCCC AGCCGCCAT GGCCCAGTC CAGTCGACAG  
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2351 GTGGAGGGCG TTACGGCGGA GTGGCTCTG CGGGTGGCGG AAGTGGACTC  
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2451 CGCGTGGCG CATAGGCTGG CGCGGGCTCT GTGGTGGTT CTGGTGGCGG  
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2501 CTCTGAGGGT GCGGGCTCTG AGGGTGGCGG TTCTGAGGGT GGCGGCTCTG  
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2551 AGGGTGGCGG TTCCGGTGGC GGCTCCGGTT CGGGTGATT TGATTATGAA  
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2601 AAAATGGCAA ACGCTAATAA GGGGGCTATG ACCGAAAATG CCGATGAAAA  
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2651 CGCGCTACAG TGTGACGCTA AAGGCAAACT TGATTCGTC GCTACTGATT  
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2701 ACGGTGGCTC TATGCTGTTG TTCAATTGGT ACGTTTCCGG CCTTGTCTAAT  
TGCCACGAC ATAGCTACCA AAGTAACAC TGCAAGGGC GGAACGATTAA  
2751 GGTAAATGGTG CTACTGGTGA TTTTGTGGC TCTAATTCCC AAATGGCTCA  
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-Replacement Sheet-  
11/43

2801 AGTCGGTGAC GGTGATAATT CACCTTAAT GAATAATTTC CGTCAATATT  
2851 TCAGCCACTG CCACATTAA GTGGAATTAA CTTATTAAGG GCAGTTTAA  
2901 TACCTCTT GGCTCAGTCG GTTGAATGTC GCCCTTATGT CTTGGCGCT  
ATGGAAGAAA CGGAGTCAGC CAACCTACAG CGGAAATACA GAAACCGCGA  
2951 GTGAAACCAT ATGAATTTC TATTGATTGT GACAAAATAA ACTTATTCCG  
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2951 TGTTGCTTT CGCTTCTT TATACTTGC CACCTTTATG TATGTTTT  
ACCACAGAAA CGCAAAAGAA ATATACAACGG GTGGAATAC ATACATAAAA  
3001 CGACGTTGC TAACATACTG CGTAATAGG AGTCTTAATA AGAATTCACT  
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3051 GGCGCTCGGT TTACAAACGTC GTGACTGGGA AAACCGTGGC GTTACCCAAC  
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3101 TTAATCGCT TGACGCCACAT CCCCTTTCG CCAGCTGGG TAATAGCAGA  
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3151 GAGGGCCCGA CGGATGCCG TTCCCAACAG TTGCGCAGCC TGAATGGCGA  
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3201 ATGGGCGCTG ATGGGGTATT TTCTCCCTAC GCATCTGTGC GGTTATTTCAC  
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3351 GCGCTAGCGG CCGCTCTT CGCTTCTTC CCTTCCCTTC TCGCACCGT  
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3451 GATTAGTGC TTACGGCACT CTGACGCCA AAAAACCTTG ATTGGGTGAT  
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3501 GTGGCACGTA GTGGGCCATC GCGCTGTAG ACGGTTTTTC GTCCCTTGAC  
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3551 GTTCGAGTC ACCTTCTTA ATAGTGGACT CTGTTCCAA ATGGAAACAA  
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3601 TACTCAACCC TATGCCGGC TATTCTTTG ATTATAAGG GATTTGGCG  
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3651 ATTGGGCCT ATTGGTTAA AAATGAGCTG ATTAAACAAA ATTAAACCG  
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3701 GAATTAAAC AAAATATTA CGTTTACAT TTATGGTG AGTCTCAGTA  
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3751 CAATCTGTC TGATGCCGCA TAGTTAACCC AGCCCCGACA CCCGCCAAC  
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3801 CCCGTGACG CGCCCTGACG GGCTTGTCTG CTCCGGCAT CGCCTACAG  
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3851 ACAAGCTGZ ACCGTCCTGG GGAGCTGCA GTGTCAGAGG TTTTCAACCGT  
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3901 CATCACCGAA ACGCCGGA GTAGTGGCTT TGCGCGCT

Fig. 7b

Nucleotide Sequences pXP14 (SEQ ID No: 112)

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51 ATAATGGTTT CTTAGACGTC AGGTGGCACT TTTCGGGAA ATGTGCGGG  
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101 AACCCCTATT TGTTTATT TTCTAAATACA TTCAAATATG TATCCGCTCA  
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151 TGAGACAATA ACCCTGATAA ATGCTTCAT AATATTGAA AAGGAAGAGT  
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201 ATGAGTATTAC AACATTCCG TGTCGCCCTT ATTCCCTTTT TTGCGGATT  
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251 TTGCGCTTCT GTTTCGCTC ACCAGAAC GCTGTTGAAA GTAAAAGATG  
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301 CTGAAGATCA GTTGGGTGCT CGAGTGGGTT ACATCGAACT GGATCTAAC  
GACTCTGTTAC CAACCCACGA GCTCACCCAA TGAGCTTGA CCTAGAGTTG  
351 AGCGGAAGA TCTCTGAGAG TTTTCGCGCC GAAGAACGTT TTCCAATGAT  
TCGCCATTCT AGGAACCTCTC AAAAGCGGGG CTTCTTGCAA AAGGTTACTA  
401 GAGCACTTTT AAAGTCTGC TATGTCGGCG GGTTATTATCC CGTATTGACG  
CTCGTGAAAA TTTCAGACG ATACACCCGG CCATAATAGG GCATAACTGC  
451 CGGGCAAGA GCAACTCGGT CGGCCATCAT ACTATTCTCA GAATGACTTG  
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501 GTTGGACTT CACCACTGAC AGAAAAGCAT CTTACGGATG GCATGACAGT  
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551 AAGAGAATTA TGCAGTCTG CCATAACCTC GAGTGTATAAC CTGCGGCCA  
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601 ACTTACTCTT GACAACGATC GGAGGACCGA AGGAGCTAAC CGCTTTTTG  
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651 CACAACATGG GGGATCATGA AACTCGCTT GATGTTGGG AACCGGAGCT  
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701 GAATGAGGCC ATACCAAAACG AGCAGCGTGA CACCCAGATG CCTGTAGCAA  
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751 TGGCAACAAAC GTTGCACAACTA CTATTAACCTG GCGAACTACT TACTCTAGCT  
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801 TCCCGGCAAC AATTAAATAGA TCGGTGAGG GCGGATAAAAG TTGCGAGGAC  
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851 ACTTCTGCGC TCGGCCCTC CGGCTGGCTG GTTTATTGCT GATAAAATCTG  
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901 GAGCCGGTGA CGCTGGGTCT CGCGGTATGA TTGCGACACT GGGGCCAGAT  
CTCGGGCAACT CGCACCCAGA GCGCGATAGT AACGTCGTGA CCCCGGCTA  
951 GGTAAGCCCTT CCGGTATCGT AGTTATCTAC ACGACGGGGA GTCAAGGCAAC  
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1001 TATGGATGAA CGAAATAGAC AGATCGCTGA GATAGGTGCC TCACGTGATTA  
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1051 AGCATTGGTA ACTGTCAGAC CAAGTTACT CATATATACT TTAGATTGAT  
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1101 TAAAAACTTC ATTTTAAATT TAAAGGATC TAGGTGAAGA TCCTTTTGA  
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1151 TAATCTCATG ACCAAAATCC CTTAACGTGA GTTTCGTTT CACTGAGCGT  
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1201 CAGACCCCGT AGAAAAGATC AAAGGATCTT CTTGAGATCC TTTTTTCTG  
GTCTGGGCA TCTTTCTAG TTTCTAGAA GAACCTCTAGG AAAAAAAGAC

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1251 CGCGTAAATCT GCTGCTTGCA AACAAAAAAA CCACCGCTAC CAGCGGTGGT  
GCCCATTTAGA CGACGAACGT TTGTTTTTTT GGTTGGCGATG TGCGCACCA  
1301 TTGTTTGGC GATCAAAGGC TACCAACTCT TTTCGGAG AAAAGGCTTC CATTGACCGA  
ACAAACAGGC CTAGTCTTCG ATGGTTGAGA AAAAGGCTTC CATTGACCGA  
1351 TCAGCAGAGC GCAGATACCA AATACTGTCC TTCTAGTGTA GCCGTAGTTA  
AGTCGTCCTCG CGTCATGTT TTATGACAGG AAGATCACAT CGGCATCAAT  
1401 GGCCACCAACT TCAAGAACCTC TGAGCACCG CCTACATACC TCGCTCTGCT  
CCGGTGGTGA AGTTCTTGAG ACATCGTGG GGATGTATGG AGCGAGACGA  
1451 AATCCTGTTA CCAGTGGCTG CTGCCAGTGG CGATAAGTCG TGTCTACCG  
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1601 ACTGAGATAC CTACAGCGT AGCTATGAGA AAGCGCCACG CTTCGCGAAG  
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1701 CGCACGAGGG AGCTTCCAGG GGAAACGCC TGGTATCTT ATAGTCCGT  
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1951 GCCGCAGCGC AACGACCGAG CGCACCGAGT CAGTGAGCGA GGAAGCGGAA  
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2001 GAGCGCCCAA TAGCRAAACCC GCCTCTCCC CGCGGTGTTGC CGATTCTTA  
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2101 AGCGAAATTAA TGTGAGTTAG CTCACTCATC AGGCACCCCA GGCTTACAC  
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2301 GCTGGATTGT TATTACTCGC GGCGGAGGCC GCCATGGCCC AGGTGCAGCT  
CGACCTAACAA ATAATGAGCG CGGGTCCGGC CGGTACCGGG TCCACCGTCA  
2351 GCAGGTCGGC CTCGAGATCA AACGGGGCGGC CGCAGGTGCGG CCGGTGCGGT  
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2401 ATCCAGATCC GCTGGAACCG CGTGGGGCCG CAAGCGCTTG GAGCCACCG  
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2451 CAGTTCGAAA ATAATAAGG ATCCGAATTC ACTGGCGTCG TTGTTACAAAC  
GTCAAGCTTT TTATTATTCC TAGGCTTAAG TGACCGGCAG CAAATGTTG  
2501 GTCGTGACTG GGAAAACCTT GGCGTTACCC AACTTAATCG CTTGCGAGCA

-Replacement Sheet-  
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CAGCACTGAC CCTTTGGGA CGCGAATGGG TTGAATTAGC GGAACGTCGT  
2551 CATCCCCCTT TCGCCAGCTG GCGTAATAGC GAAGAGGCC GCACCGATCG  
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2651 ATTTTCTCT TACGCATCTG TGCGGTATTT CACACCGCAT ACGTCAAAGC  
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2701 AACCATAGTA CGCGCCCTGT AGCGGCCGAT TAAGCCCGC GGGTGTGGT  
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2751 GTTACGGCGA CGGTGACCGC TACACTTGCC AGGCCCTAG CCCCCCGTCC  
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3001 TTAATAGTGG ACTCTTGTG CAAACTGGAA CAATACTCAA CCTATCTCG  
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3051 GCCTATTCTT TTGATTATAG AGGGATTTC CCGATTTCGG CCTATTGGTT  
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3101 AAAAATGAG CTGATTAAAC AAAAATTAA CGCGAATTTC AACAAAATAT  
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3151 TAACGTTTAC AATTTTATGG TGCACTCTCA GTACAATCTG CTCTGATGCC  
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3201 GCATAGTTAA GCCAGCCCC ACACCCGCCA ACACCCGCTG ACGCCCTG  
CGTATCAATT CGGTGGGGGC TGTGGGGCGT TGTGGGGCAC TGCGCGGGAC  
3251 ACGGGCTTGT CTGCTCCCGG CATCCGCTTA CAGACAAGCT GTGACCGTCT  
TGGCCGAACAA GACGAGGGCC GTAGGGCAAT GTCTGTTGCA CACTGGCAGA  
3301 CGGGGAGCTG CATGTGTCAAG AGGTTTCAC CGTCATCACC GAAACGCGCG  
GCCCTCGAC GTACACAGTC TCCAAAAGTG GCAGTAGTGG CTTTGCACGC  
3351 A  
T

Fig. 8b

-Replacement Sheet-  
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**cDNA primers**

VLK-c	CTGGATGGTGGGAAAGATGGA (SEQ ID No:113)
VLL-c	TCAGAGGAAGGAAACAGGGT (SEQ ID No:114)
IgG1-c	CTTACAACCACAATCCCTGGGCACAATT (SEQ ID No:115)
IgG2a-c	CTTTGTGGGCCCTCTGGGCTCAAT (SEQ ID No:116)
IgG2b	TGAAATGGGCCGCTGGGCTCAAG (SEQ ID No:117)
IgG3-c	GGGCTTGGGTATTCTAGGCTCGAT (SEQ ID No:118)

**VH forward primers without restriction sites**

M-VH1	GAGGTGCAGCTTCAGGAGTCAGG (SEQ ID No:119)
M-VH2	CAGGTGCAGCTGAAGGAGTCAGG (SEQ ID No:120)
M-VH3	GAGGTCCAGCTGCAACAGCTGG (SEQ ID No:121)
M-VH4	GAGGTTCAGCTGCAGCAGTCAGG (SEQ ID No:122)
M-VH5	CAGGTCACACTGCAGCAGCTGG (SEQ ID No:123)
M-VH6	CAGGTTCACTGCAGCAGTCAGG (SEQ ID No:124)
M-VH7	GAGGTGAAGCTGGTGGAGTCAGG (SEQ ID No:125)
M-VH8	GAGGTGAAGCTGGTGGAAATCTGG (SEQ ID No:126)
M-VH9	GAGGTTCACTGCAGCAGTCAGG (SEQ ID No:127)

**VH backward primers without restriction sites**

M-JH1	TGAGGAGACGGTGACCGTGGTCCC (SEQ ID No:128)
M-JH2	TGAGGAGACTGTGAGAGTGGTGCC (SEQ ID No:129)
M-JH3	TGCAGAGACAGTGACCAAGAGTCCC (SEQ ID No:130)
M-JH4	TGAGGAGACGGTGACTGAGGTTCC (SEQ ID No:131)

**VL forward primer without restriction sites**

M-VK1	GACATTGATGACACAGTCCTCC (SEQ ID No:132)
M-VK2	GATGTTGATGACCCAAACTCC (SEQ ID No:133)
M-VK3	GATATCCAGATGACACAGACTCC (SEQ ID No:134)
M-VK4	CAAATTGTTCTCACCCAGTCCTCC (SEQ ID No:135)
M-VL1	CAGGCTGTTGTGACTCAGGAATC (SEQ ID No:136)

**VL backward primers without restriction sites**

M-JK1	TTTATTTCAGCTTGGTCCCTCC (SEQ ID No:137)
M-JK2	TTTTATTTCAGCTTGGTCCCCCC (SEQ ID No:138)
M-JK3	TTTCAGCTCAGCTTGGTCCAGC (SEQ ID No:139)
M-JL1	ACCTAGGACAGTGACCTTGGTCC (SEQ ID No:140)

**VH forward primers with restriction sites**

MVH1 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCGAGGTGCAGCTTCAGGAGTCAGG (SEQ ID No:141)
MVH2 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCCCAGGTGCAGCTGAAGGAGTCAGG (SEQ ID No:142)
MVH3 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCGAGGTCCAGCTGCAACAGTCTGG (SEQ ID No:143)
MVH4 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCGAGGTTCAGCTGCAGCAGTCTGG (SEQ ID No:144)
MVH5 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCCCAGGTCAAUTGCAGCAGCTGG (SEQ ID No:145)
MVH6 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCCCAGGTTCAGCTGCAGCAGTCTGG (SEQ ID No:146)
MVH7 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCGAGGTGAAGCTGGTGGAGTCTGG (SEQ ID No:147)
MVH8 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCGAGGTGAAGCTGGTGAAGTCTGG (SEQ ID No:148)
MVH9 Sfil	GTCCTCGCAACTGCGGCCAGCCGGCATGGCGAGGTTCAGCTTCAGCAGTCTGG (SEQ ID No:149)

**VH backward primers with restriction sites**

MJH1 Sall	GAGTCATTCTCGTGTGACACAGTGACCGTGAACGGTCCC (SEQ ID No:150)
MJH2 Sall	GAGTCATTCTCGTGTGACACTGTGAGAGTGGTGC (SEQ ID No:151)
MJH3 Sall	GAGTCATTCTCGTGTGACACAGTGACCAAGATCCC (SEQ ID No:152)
MJH4 Sall	GAGTCATTCTCGTGTGACACAGTGACTGAGGTCCC (SEQ ID No:153)

**VL forward primers with restriction sites**

MVK1 ApaL1	TGAGCACACAGTCACCTCGACATTGTGATGACACAGTCTCC (SEQ ID No:154)
MVK2 ApaL1	TGAGCACACAGTCACCTCGATGTGATGACCCAACTCC (SEQ ID No:155)
MVK3 ApaL1	TGAGCACACAGTCACCTCGATATCCAGATGACACAGTCTCC (SEQ ID No:156)
MVK4 ApaL1	TGAGCACACAGTCACCTCCAAATTGTTCTACCCAGTCTCC (SEQ ID No:157)
MVL1 ApaL1	TGAGCACACAGTCACCTCCAGGTGACTGAGGTCCC (SEQ ID No:158)

**VL backward primers with restriction sites**

M-JK1 Not1	GAGTCATTCTCGACTTGGCCGCTTGATTTCCAGCTTGGTGCCTCC (SEQ ID No:159)
M-JK2 Not1	GAGTCATTCTCGACTTGGCCGCTTTATTCAGCTTGGTCCCCCCC (SEQ ID No:160)
M-JK3 Not1	GAGTCATTCTCGACTTGGCCGCTTCAGCTCCAGCTTGGTCCCAGC (SEQ ID No:161)
M-JL1 Not1	GAGTCATTCTCGACTTGGCCGACCTAGGACAGTGCACCTGGTCCC (SEQ ID No:162)

Fig. 9